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Key indicators

Single-crystal X-ray study T = 142 K Mean σ (C–C) = 0.002 Å R factor = 0.036 wR factor = 0.099 Data-to-parameter ratio = 13.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

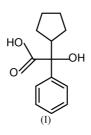
2-Cyclopentyl-2-hydroxy-2-phenylacetic acid, a pseudosymmetric crystal structure with space group $Pna2_1$ and Z' = 2

The title compound, $C_{13}H_{16}O_3$, crystallizes in space group $Pna2_1$ with Z = 8. The two independent molecules are related by pseudosymmetry elements. The structure can be derived from the higher symmetric space group *Pnam* by translating the *x* coordinates of molecule 2 by -1/4 with respect to those of molecule 1.

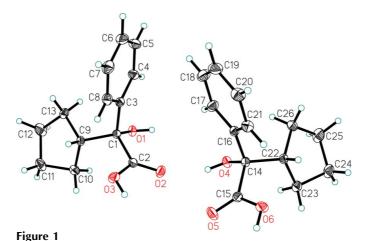
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Comment

The title compound, (I), was prepared as a starting material for the synthesis of possible anticholinergic agents.



The structure contains two crystallographically independent molecules (Fig. 1). The dimensions of both molecules are very similar. The maximum differences for corresponding bond distances and angles are 0.021 Å for the C2–O2 and C15–O5 bonds, and 0.9° for the C6–C7–C8 and C19–C20–C21 angles. Corresponding torsion angles in the two molecules agree within 2°. The phenyl groups are planar. The cyclopentyl rings have approximately envelope conformations: a C10-envelope for molecule 1 and a C23-envelope for molecule 2. The molecular conformation of (I) is very similar to the



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The structures of the two independent molecules of (I), shown with 50% probability displacement ellipsoids.

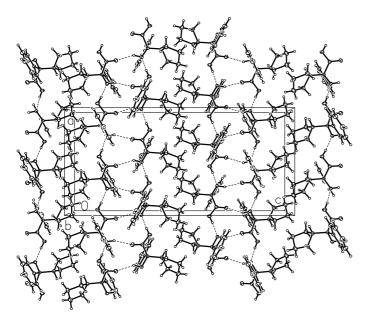


Figure 2 The crystal packing of (I), viewed down b.

conformation of a 1,1-dimethylpyrrolidinium ester of (I), reported by Guy & Hamor (1973). Only the conformation of the cyclopentyl group is considerably different in that compound.

The molecules are connected by $O_{carboxyl} - H \cdots O_{hydroxyl}$ hydrogen bonds to form chains in the crystallographic a direction (Table 1 and Fig. 2). One chain consists only of molecule 1, while the next chain consists only of molecule 2. The molecules within each chain are related by crystallographic a-glide-plane symmetry. Two neighboring chains are linked by $O_{hydroxyl}$ – $H \cdots O_{keto}$ hydrogen bonds to form double chains. These double chains are further stabilized by weak intermolecular $C-H \cdot \cdot \pi$ interactions between phenyl groups: $C5-H5\cdots Cg_2$ (where Cg_2 is the centroid of the phenyl ring of molecule 2) and C18-H18···C5C6 (where C5C6 is the midpoint of the C5–C6 bond). Weak intermolecular C_{phenyl}- $H \cdots O_{hvdroxvl}$ interactions, with $H \cdots O$ distances of 2.67 and 2.69 Å, connect neighboring chains in the b direction.

The ADDSYM option of program *PLATON* (Spek, 1998) revealed an approximate inversion center at position (3/8, 1/2, 1/2)0), indicating the existence of pseudosymmetry. The coordinates of molecule 2 are related to those of molecule 1 by the approximate relations: $x_2 = 1/4 + x_1$, $y_2 = y_1$ and $z_2 = 1/2 - z_1$. This corresponds to a non-crystallographic pseudo-glide-plane perpendicular to the c axis, with a translation of a/4. Molecule 2 is related to symmetry equivalents of molecule 1 by a pseudo-inversion center at (1/8, 0, 0) (which is a position equivalent to the pseudo-inversion center detected by *PLATON*), a pseudo-twofold screw axis along (x, 1/4, 1/4), with a translation vector of 3a/4, and a pseudo-twofold screw axis along (3/8, y, 0), with a translation vector of b/2. These pseudosymmetry operations are local symmetry elements which relate molecule 1 to molecule 2.

The structure of (I) is closely related to the higher symmetric space group *Pnam* (non-standard setting of *Pnma*). If the x coordinates of the atoms of molecule 2 are shifted by -1/4, then the local pseudo-glide-plane in (I) is replaced by the mirror plane at z = 1/4 of space group *Pnam*. A search of the Cambridge Structural Database (Allen, 2002) for crystal structures with space group Pna2₁, reported with two independent molecules, revealed a number of other structures with pseudosymmetry relations. Examples are anthranilic acid (Brown & Ehrenberg, 1985), pyrazole (Larsen et al., 1970; la Cour & Rasmussen, 1973), dimethylaminesulfur dioxide (Phillips et al., 1996) and 3-amino-4-nitrofurazan (Batsanov & Struchkov, 1985). All these structures can be derived from the higher symmetric space group Pnma by translating some molecules with respect to their neighbors.

Experimental

The methyl ester of the title compound was prepared following a procedure described by Smith et al. (1949) and Adlerová et al. (1964). 60 ml of an ethereal Grignard solution made from magnesium turnings (2.7 g, 111 mmol) and bromocyclopentane (16 ml, 150 mmol) was added dropwise to an ice-cooled solution of 2-oxo-2-phenylacetic acid methyl ester (7 g, 42.6 mmol) in 70 ml of dry diethyl ether. Aqueous work-up and flash chromatography (silica gel, petroleum ether/ethyl acetate, 20:1) afforded 3.3 g (33%) of 2-cyclopentyl-2hydroxy-2-phenylacetic acid methyl ester. The title compound, (I), was obtained by hydrolysis of this ester with lithium hydroxide monohydrate in a mixture of THF and water (1:1). Single crystals of (I) were recrystallized from diethyl ether.

Crystal data

$\begin{array}{l} C_{13}H_{16}O_{3} \\ M_{r} = 220.26 \\ Orthorhombic, Pna2_{1} \\ a = 11.1981 \ (14) \ \text{\AA} \\ b = 8.4864 \ (15) \ \text{\AA} \\ c = 23.853 \ (3) \ \text{\AA} \\ V = 2266.8 \ (6) \ \text{\AA}^{3} \\ Z = 8 \\ D_{x} = 1.291 \ \text{Mg m}^{-3} \end{array}$	Mo K α radiation Cell parameters from 194 reflections $\theta = 3-23^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ T = 142 (2) K Rod, colorless $0.50 \times 0.45 \times 0.40 \text{ mm}$	
Data collection		
Siemens SMART 1K CCD diffractometer ω scans Absorption correction: multi-scan (<i>SADABS</i> , Sheldrick, 2000) $T_{min} = 0.935$, $T_{max} = 0.964$ 37 645 measured reflections	3878 independent reflections 3597 reflections with $I > 2\sigma(I)$ $R_{int} = 0.023$ $\theta_{max} = 32.0^{\circ}$ $h = -16 \rightarrow 16$ $k = -11 \rightarrow 10$ $l = -34 \rightarrow 35$	
Refinement		
Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.036$ $wR(F^2) = 0.099$ S = 1.08 3878 reflections 294 parameters H-atom parameters constrained	$\begin{split} w &= 1/[\sigma^2(F_o{}^2) + (0.06P)^2 \\ &+ 0.35P] \\ \text{where } P &= (F_o{}^2 + 2F_c{}^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.001 \\ \Delta\rho_{\text{max}} &= 0.39 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.23 \text{ e } \text{\AA}^{-3} \\ \text{Extinction correction: } SHELXL97 \\ \text{Extinction coefficient: } 0.0043 (10) \end{split}$	

Table 1

Hydrogen-bonding geometry (Å, °).

 Cg_2 is the centroid of the phenyl ring of molecule 2 and C5C6 is the midpoint of the C5-C6 bond.

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$O1-H01\cdots O5^i$	0.84	2.00	2.794 (2)	156
O3-H03···O1 ⁱⁱ	0.84	1.83	2.664 (2)	173
$O4-H04\cdots O2$	0.84	2.00	2.762 (2)	151
O6−H06···O4 ⁱⁱ	0.84	1.78	2.621 (2)	174
$C5-H5\cdots Cg_2^{iii}$	0.95	2.82	3.521 (2)	132
C18-H18···C5C6	0.95	2.73	3.582 (2)	150
$C6-H6\cdots O1^{iv}$	0.95	2.67	3.587 (2)	163
C19−H19····O4 ^{iv}	0.95	2.69	3.601 (2)	160

x, y - 1, z.

Equivalent reflections, including Friedel opposites, were merged, since anomalous dispersion effects are negligible. The systematic absences showed the space group to be either $Pna2_1$ or Pnam (non-standard setting of Pnma). The almost centrosymmetric distribution of the *E* values and the value of Z (Z = 8) are indications of the latter space group. All attempts to determine the structure in space group Pnam, however, failed. Consequently, the structure was determined in the non-centrosymmetric space group $Pna2_1$. The resulting structure can not be transformed to the centrosymmetric space group Pnam. The H atoms were located in a difference Fourier synthesis. They were refined with fixed individual displacement parameters $[U_{iso}(H) = 1.2U_{eq}(C), U_{iso}(H) = 1.5U_{eq}(C_{methyl})$ and $U_{iso}(H) = 1.5U_{eq}(O)]$ using a riding model, with fixed distances: H-O = 0.84 Å, H-C(phenyl) = 0.95 Å, H-C(secondary) = 0.99 Å and H-

C(primary) = 1.00 Å. The hydroxyl and carboxyl H atoms were allowed to rotate about the O-C bonds.

Data collection: *SMART* (Siemens, 1995); cell refinement: *SMART*; data reduction: *SAINT* (Siemens, 1995); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 1996); software used to prepare material for publication: *SHELXL*97.

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